AMENDMENT UNDER 37 C.F.R. § 1.111 Attorney Docket No.: Q83564

Application No.: 10/511,098

## **AMENDMENTS TO THE CLAIMS**

This listing of claims will replace all prior versions and listings of claims in the application:

## I. LISTING OF CLAIMS:

- 1-32. (cancelled)
- 33. (currently amended) An expression vector,

  which comprises comprising: (a) a first coding region encoding PPIasea peptidyl
  prolyl cis-trans isomerase (PPIase) having molecular chaperone activity, and
- (b) a region having at least one restriction enzyme site in which a second coding region encoding a desired protein can be inserted.
- 34. (previously presented): The expression vector according to claim 33, wherein the first coding region is operatively linked to a promoter, and the restriction enzyme site is in the same reading frame as the first coding region, and is downstream of the first coding region.
- 35. (currently amended): The expression vector according to claim 33, which has a region being between athe first coding region and athe region having at least one restriction enzyme site in which a second coding region can be inserted, and is translated in the same reading frame to be a protease digestion site in the same reading frame as the first and second coding regions.
- 36. (previously presented): An expression vector, wherein a second coding region encoding a desired protein is inserted into the expression vector according to claim 33.
- 37. (previously presented): The expression vector according to claim 33,

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- wherein the PPIase having molecular chaperone activity is FKBP-type PPIase.
- 38. (previously presented): The expression vector according to claim 33, wherein the PPIase having molecular chaperone activity is cyclophilin-type PPIase.
- 39. (previously presented): The expression vector according to claim 33, wherein the PPIase having molecular chaperone activity is parvulin-type PPIase.
- 40. (previously presented): The expression vector according to claim 37, wherein the FKBP-type PPIase is archaebacterial FKBP-type PPIase.
- 41. (previously presented) The expression vector according to claim 40, wherein the archaebacterial FKBP-type PPIase is short type FKBP-type PPIase.
- 42. (previously presented): The expression vector according to claim 33, wherein the PPIase having molecular chaperone activity comprises an IF domain and/or a C-terminal domain of archaebacterial FKBP-type PPIase.
- 43. (previously presented): The expression vector according to claim 37, wherein the FKBP-type PPIase is trigger factor-type PPIase.
- 44. (previously presented): The expression vector according to claim 33, wherein the PPIase having molecular chaperone activity comprises a N-terminal domain and/or a C-terminal domain of trigger factor-type PPIase.
- 45. (previously presented): The expression vector according to claim 37, wherein the FKBP-type PPIase is FkpA-type PPIase.
- 46. (previously presented): The expression vector according to claim 33, wherein the PPIase having molecular chaperone activity comprises a N-terminal domain of FkpA-type PPIase.

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47. (previously presented): The expression vector according to claim 37, wherein the FKBP-type PPIase is FKBP52-type PPIase.

- 48. (previously presented): The expression vector according to claim 33, wherein the PPIase having molecular chaperone activity comprises a C-terminal domain of FKBP52-type PPIase.
- 49. (previously presented): The expression vector according to claim 38, wherein the cyclophilin-type PPIase is CyP40-type PPIase.
- 50. (previously presented): The expression vector according to claim 33, wherein the PPIase having molecular chaperone activity comprises a C-terminal domain of CyP40-type PPIase.
- 51. (previously presented): The expression vector according to claim 39, wherein the parvulin-type PPIase is SurA-type PPIase.
- 52. (previously presented): The expression vector according to claim 33, wherein the PPIase having molecular chaperone activity comprises a N-terminal domain of SurA-type PPIase.
- 53. (previously presented): The expression vector according to claim 36, wherein the second coding region has a nucleotide sequence encoding a monoclonal antibody.
- 54. (previously presented): The expression vector according to claim 36, wherein the second coding region has a nucleotide sequence encoding a membrane protein.
- 55. (previously presented): A host,which contains the expression vector according to claim 33.

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- 56. (previously presented): The host according to claim 55, which is Escherichia coli.
- 57. (previously presented) A fused protein,
  which comprises PPIase having molecular chaperone activity and a desired protein.
- 58. (previously presented): The fused protein according to claim 57, which comprises a protease digestion site between PPIase having molecular chaperone activity and a desired protein.
- 59. (currently amended): A process for producing a fused protein comprising PPIase having molecular chaperone activity and a desired protein,

which comprises making the expression vector according to claim 36, comprising culturing a host cell transformed with the expression vector of claim 33 to express the fused protein.

60. (currently amended): The process for producing a fused protein according to claim 59,

which comprises culturing the a host containing the expression vector under condition of expression of the expression vector, and making express expressing the fused protein in a cytoplasm.

61. (currently amended): The process for producing a fused protein according to claim 59,

which comprises providing a region being transcribed and translated to be a signal sequence at a 5' terminus of athe first coding region or a 5'3' terminus of athe second coding region of the expression vector, and culturing a host containing the expression

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vector under condition of expression of the expression vector to express the fused protein in athe periplasm or a medium.

62. (currently amended): The process for producing a fused protein according to claim 59,

which comprises making the expression vector culturing a host cell transformed with the expression vector to express the fused protein in a cell-free translation system.

63. (currently amended): The process for producing a fused protein according to claim 59,

wherein the fused protein is adsorbed on a carrier harboring macrolide, cyclosporin, juglone, or its analogous a compound inhibiting which inhibits PPIase activity, and then the wherein said carrier is recovered and the fused protein is recovered from the carrier.

64. (currently amended): A process for producing a desired protein,

which comprises digesting the fused protein comprising athe protease digestion site obtained by the process according to claim 59, with a protease digesting a protease digestion site.